

AAGGTGTTTGAGCTTGTGGGTGAGCCCTCCATATACTGCACCAGCAATGACGATCAAGTG
GGCATCTGGAGCGGCCCGGCCCTCAGTGCATTATACCTAACAAATGCACGCCTCCAAAT
GTGGAAAATGGAATATTGGTATCTGACAACAGAGCTTATTTTCCTTAAATGAAGTTGTG
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GCTGGAATGGAAAGCCTTTGGAATAGCAGTGTTCAGTGTGTGAACAAATCTTTTGTCCA
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ATTGGAGAGAGCACCATCCGCTGCACAAGTGACCCTCAAGGGAATGGGGTTTGGAGCAGC
CCTGCCCCCTCGCTGTGGAATTATCACCATCACCATCACTAAAGATCT

Fig. 9B (cont.)

Fig. 10A

SEQ. ID NO:17

ATA TAC GAA TTC TGG TTG AGT CCA AAT ATG GTC CC

Fig. 10B

SEO, ID NO:18

ACA GTG AGA TCT TTA TCA TTT ACC CGG AGA CAG GGA G

Fig. 11A

SEQ. ID NO:19

	M T V A R P S V P A A L P																			
L	L	G	E	L	P	R	L	T	L	L	V	L	L	C	L	P	A	V	W	G
D	C	G	L	P	P	D	V	P	N	A	Q	P	A	L	E	G	R	T	S	F
P	E	D	T	V	I	T	Y	K	C	E	E	S	F	V	K	I	P	G	E	K
D	S	V	I	C	L	K	G	S	Q	W	S	D	I	E	E	F	C	N	R	S
C	E	V	P	T	R	L	N	S	A	S	L	K	Q	P	Y	I	T	Q	N	Y
F	P	V	G	T	V	V	E	Y	E	C	R	P	G	Y	R	R	E	P	S	L
S	P	K	L	T	C	L	Q	N	L	K	W	S	T	A	V	E	F	C	K	K
K	S	C	P	N	P	G	E	I	R	N	G	Q	I	D	V	P	G	G	I	L
F	G	A	T	I	S	F	S	C	N	T	G	Y	K	L	F	G	S	T	S	S
F	C	L	I	S	G	S	S	V	Q	W	S	D	P	L	P	E	C	R	E	I
Y	C	P	A	P	P	Q	I	D	N	G	I	I	Q	G	E	R	D	H	Y	G
Y	R	Q	S	V	T	Y	A	C	N	K	G	F	T	M	I	G	E	H	S	I
Y	C	T	V	N	N	D	E	G	E	W	S	G	P	P	P	E	C			
S	S	P	N	K	C	T	P	P	N	V	E	N	G	I	L	V	S	D	N	